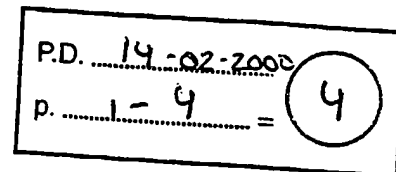```
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DT   24-SEP-2001 (Rel. 69, Last updated, Version 8)
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DE   Homo sapiens chromosome 11 clone RP11-583F24 map 11, *** SEQUENCING IN
DE   PROGRESS ***, 2 ordered pieces.
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KW   HTG; HTGS_ACTIVEFIN; HTGS_FULLTOP; HTGS_PHASE2.
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RL   Unpublished.
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RA   Raymond C., Riley R., Rothman D., Roy A., Santos R., Severy P., Spencer B.,
RA   Stange-Thomann N., Stojanovic N., Subramanian A., Talamas J., Tesfaye S.,
RA   Theodore J., Tirrell A., Vassiliev H., Viel R., Vo A., Wu X., Wyman D.,
RA   Ye W.J., Zimmer A., Zody M.;
RT   ;
RL   Submitted (08-FEB-2000) to the EMBL/GenBank/DDBJ databases.
RL   Whitehead Institute/MIT Center for Genome Research, 320 Charles Street,
RL   Cambridge, MA 02141, USA
XX
CC   On Sep 20, 2001 this sequence version replaced gi:13559563.
CC   All repeats were identified using RepeatMasker:
CC   Smit, A.F.A. & Green, P. (1996-1997)
CC   http://ftp.genome.washington.edu/RM/RepeatMasker.html
CC   -------------- Genome Center
CC   Center: Whitehead Institute/ MIT Center for Genome Research
CC   Center code: WIBR
CC   Web site: http://www-seq.wi.mit.edu
CC   Contact: sequence_submissions@genome.wi.mit.edu
CC   -------------- Project Information
CC   Center project name: L6254
CC   Center clone name: 583_F_24
CC   --------------.
CC   * NOTE: This is a 'working draft' sequence. It currently
CC   * consists of 2 contigs. Gaps between the contigs
CC   * are represented as runs of N. The order of the pieces
CC   * is believed to be correct as given, however the sizes
CC   * of the gaps between them are based on estimates that have
CC   * provided by the submittor.
CC   * This sequence will be replaced
CC   * by the finished sequence as soon as it is available and
CC   * the accession number will be preserved.
CC   *   1   81101: contig of 81101 bp in length
CC   * 81102 81201: gap of   100 bp
CC   * 81202   164531: contig of 83330 bp in length.
XX
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XP-002192686

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